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Publication Title	Authors	Journal (Pub date)	Volume/Page	PubMed Li
A canonical promoter organization of the transcription machinery and its regulators in the Saccharom	Venters, Bryan J; Pugh, B Franklin	Genome Res (2009 Mar)	19 / 360-71	PubMed Citat
A comprehensive and high-resolution genome-wide response of p53 to stress.	Chang, Gue Su; Chen, Xiangyun Amy; Park, Bongsoo; Rhee, Ho Sung; Li, Pingxin; Han, Kang Hoo; Mishra, Tejaswini; Chan-Salis, Ka Yim; Li, Yunfei; Hardison, Ross C; Wang, Yanming; Pugh, B Franklin	Cell Rep (2014 Jul 24)	8 / 514-27	PubMed Citat
A comprehensive genomic binding map of gene and chromatin regulatory proteins in Saccharomyces.	Venters, Bryan J; Wachi, Shinichiro; Mavrich, Travis N; Andersen, Barbara E; Jena, Peony; Sinnamon, Andrew J; Jain, Priyanka; Rolleri, Noah S; Jiang, Cizhong; Hemeryck-Walsh, Christine; Pugh, B Franklin	Mol Cell (2011 Feb 18)	41 / 480-92	PubMed Citat
ChIP-exo method for identifying genomic location of DNA-binding proteins with near-single-nucleotide	Rhee, Ho Sung; Pugh, B Franklin	Curr Protoc Mol Biol (2012 Oct)	Chapter 21 / Unit 21.24	PubMed Citat
Comprehensive genome-wide protein-DNA interactions detected at single-nucleotide resolution.	Rhee, Ho Sung; Pugh, B Franklin	Cell (2011 Dec 9)	147 / 1408-19	PubMed Citat
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Sequential recruitment of SAGA and TFIID in a genomic response to DNA damage in Saccharomyces cerevi	Ghosh, Sujana; Pugh, B Franklin	Mol Cell Biol (2011 Jan)	31 / 190-202	PubMed Citat